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AAI52287
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AAH29813
AAC03878
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S cerevisiae apopt
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ALIGNMENTS

RESULT AAC49069

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18-OCT-2000 (first entry)

AAC49069 standard; DNA; 637

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Arabidopsis thaliana DNA fragment SEQ ID NO: 59811.

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          25-FEB-2000; 2000EP-0301439.
                                                   06-SEP-2000
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                                                                                                                               Arabidopsis thaliana.
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0; Mismatches 120;
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99US-0142154

99US-0142803

99US-0142803
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RESULT 3
AAC34600 :
ID AAC34600 :
XX
AC AAC34600;
XZ
DT 17-OCT-200
XX
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Best Loc
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04-CCT-1999;
05-OCT-1999;
06-OCT-1999;
07-OCT-1999;
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16-OCT-1999;
17-OCT-1999;
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18-OCT-1999;
19-OCT-1999;
             17-OCT-2000
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Local Similarity
les 195; Conserv
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                                                                                                                                                       atcgagagaatcacgagggagaaggtgaatggaacaaccgtgaacgttgggattcagccg
                                                                                                                                                                   atcgagagagtgaccagagaaaaagcgaatggaatgactgtacctgtgggaatacatcca
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                                                        standard;
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nilarity 62.3%;
Conservative
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99US-0161361
             (first entry)
                                                                                                                                     408
                                                        DNA;
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                                                           638
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Pred. No. 8e-30;
0; Mismatches 118;
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18-JUN-1999; 18-JUN-1999; 18-JUN-1999; 18-JUN-1999; 18-JUN-1999; 21-JUN-1999;	18-JUN-1999; 18-JUN-1999; 18-JUN-1999; 18-JUN-1999;	18-JUN-1999; 18-JUN-1999;	17-JUN-1999; 18-JUN-1999;	16-JUN-1999; 16-JUN-1999;	10-JUN-1999; 14-JUN-1999;	08-JUN-1999; 10-JUN-1999;	04-JUN-1999; 07-JUN-1999;	03-JUN-1999;	28-MAY-1999;	25-MAY-1999;	21-MAY-1999; 24-MAY-1999;	20-MAY-1999;	18-MAY-1999;	14-MAY-1999; 14-MAY-1999;	14-MAY-1999;	11-MAY-1999;	06-MAY-1999;	05-MAY-1999; 06-MAY-1999;	04-MAY-1999;	30-APR-1999;	23-APR-1999; 28-APR-1999;	21-APR-1999; 23-APR-1999;	19-APR-1999;	08-APR-1999;	01-APR-1999; 06-APR-1999;	29-MAR-1999;	3-MAR-19	05 MAR-1999;	в-199	25-FEB-2000;	06-SEP-2000.	EP1033405-A2.	Arabidopsis t	ic	Hybridisation as	Arabidopsis t
99US-0139463. 99US-0139463. 99US-0139750. 99US-0139763. 99US-0139763.																											99US-0125788.	99US-0123180.	99US-0121825.	2000EP-0301439.			thaliana.	ay; promoter; termination se	assay; genetic	thaliana DNA fragment SEQ ID NO: 7218.
PR 25-AUG- PR 26-AUG- PR 27-AUG- PR 27-AUG- PR 27-AUG- PR 27-AUG- PR 27-AUG-									PR 04-A	03	00	O N	N) N	1 6 1 1		K) K '			N 1 N 1													PR 01-JUL- PR 01-JUL-			PR 23-JUN PR 23-JUN	
1999; 1999; 1999; 1999; 1999;	1999;	1999;	1999;	1999;	1999;	1999;	1999;	1999;	AUG-1999; AUG-1999;	AUG-1999;	-1999;	-1999; -1999;	-1999;	-1999;	-1999; -1999:	-1999;	1999;	1999;	-1999; -1999;	-1999; -1999;	-1999;	-1999;	-1999; -1999;	1999;	.1999;	-1999;	999;	-1999;	-1999; -1999:	-1999; -1999;	-1999; -1999;	-1999; -1999;	-1999;	1999;	-1999; -1999;	-1999;
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01-SEP-1999
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                                                                                    atcgagaggattaccagagagaaggtgaattggaaccaccgtgaacgttgggattcagccg
                                atcyagagagtyaccagagaaaaagcyaatygaatgactytacctyttyggaatacatcca
                                                                                                                                                                  aaatacagcgttcgttctttacctattcgaaaagaagacgaagtaatcatagttcgcgga
                                                                                                                                                                                                                                                                                                              atgaagtacaatccaagagtgacctcttctcgcaggaagaacaggaaggctcatttcaca
                                                                                                                                                                                                                                                                                                                                                                                                      al Similarity 61.7
193; Conservative
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99US-0151303

99US-0153763

99US-0153763

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99US-0161361

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99US-0161932
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3.4e-29;
ies 120;
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  RESULT
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07 AMY-1999

11 AMY-1999

11 AMY-1999

12 AMY-1999

14 AMY-1999

14 AMY-1999

15 AMY-1999

16 AMY-1999

17 JUN-1999

20 AMY-1999

21 AMY-1999

21 AMY-1999

21 AMY-1999

22 AMY-1999

23 JUN-1999

24 AMY-1999

26 AMY-1999

27 AMY-1999

28 AMY-1999

29 JUN-1999

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05-MAR-1999
09-MAR-1999
23-MAR-1999
25-MAR-1999
29-MAR-1999
01-APR-1999
06-APR-1999
08-APR-1999
16-APR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-FEB-2000;
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|| || || || || |
358 tcgaaagttgtga
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000EP-0301439
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            QΥ
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                                                                                      Query Match 28.9%;
Best Local Similarity 62.4%;
Matches 201; Conservative
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Pred. No. 7.6e-28;
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                                   Matches
                                              Query Match
Best Local (
                                                                                                                           The present invention provides the protein and coding sequences of a number of apoptosis associated proteins from the yeast Saccharomyces cerevisiae and the fungus candida albicans. These can be used to identify treatments for fungal and yeast infections, for proliferative diseases and for apoptosis related diseases such as autoimmune diseases, ischaemia and neurodegeneration. The present sequence is one of the C. albicans
                                                                                                                                                                                                                                   Yeast and fungal nucleic acids encoding proteins involved in a pathway leading to programmed cell death, useful for treating proliferative disorders, yeast and fungal infections, or for preventing apoptosis in certain diseases -
                                                                                                                                                                                                                                                                                                                                                                (JANC
                                                                                                                                                                                                                                                                                                                                                                                                             03-JUL-2000;
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Reekmans RJ;
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                                             Sequences AAA31040-A32093 represent novel plant microsatellite sequences and associated flanking species. The sequences comprise a central core repeat sequence, especially selected from the sequences AAA32094-A32096 with left and right flanking sequences. The polynucleotide sequences can be used in the detection of DNA polymorphisms, in genome mapping, in physical mapping, in positional cloning of genes, in variety identification and in evaluation of genetic variability within and between plant tissues, populations, cultivars, species and species groups. They may also be used to design hybridization probes for oligonucleotide fingerprinting and library screening and to design primers for microsatellite-primed PCR. Microsatellite markers are useful to locate specific economically useful genes in plant genomes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1036
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     plant microsatellite sequence; core repeat sequence; detection; pro DNA polymorphism; genome mapping; physical mapping; fingerprinting; variety identification; genetic variability evaluation; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                              Page 139; 392pp;
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The present invention describes a method for monitoring differential expression of genes in a first filamentous fungal (FF) cell relative expression of the same genes in one or more second filamentous funga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Multiple gene expression; filamentous fungal cell; EST; expressed sequence tag; Fusarium veneratum; Aspergillus niger; Aspergillus oryzae; Trichoderma reesei; identification; recombination; culture condition; environmental stress; spore morphogenesis; metabolic pathway engineering; catabolic pathway engineering; ss.
                                                                                                                                                                                                                                                                                                                                                                 (NOVO)
                                                                                                                                                                               Moditoring differential expression of genes in filamentous uses fluorescence-labeled nucleic acids isolated from the c
                                                                                                             Claim 86;
                                                                                                                                                                                                                                                                  WPI; 2000-594572/56.
                                                                                                                                                                                                                                                                                                                    Berka RM,
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ous fungal
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RESULT 8
AAI52287/c
ID AAI522
XX AAI522
XX AAI522
XX Probe
XX Probe;
KW Probe;
KW geneti
XX Geneti
XX W02001
XX W02001
XX W030-AUG
PF 30-AUG
PR 04-FEH
PR 04-FEH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    engineering. Using ESTs provides several advantages over genomic or random cDNA clones including elimination of redundancy as one spot on an array equals one gene or open reading frame, and organisation of the microarrays based on function of the gene products to facilitate analysis of the results. AAF07478 to AAF11247 represents ESTs from Fusarium venenatum; AAF11248 to AAF11853 represents ESTs from Aspergillus niger; AAF11854 to AAF14878 represents ESTs from Aspergillus oryzae; and AAF14879 to AAF15337 represents ESTs from Trichoderma reesei, which are all specifically claimed in the present invention.
                                              30-JAN-2001; 2001WO-US00663
                                                                              09-AUG
                                                                                                              WO200157272-A2
                                                                                                                                                                            genetic disorder;
                                                                                                                                                                                                                      Probe #20973 used to measure gene expression in human placenta
                                                                                                                                                                                                                                                         17-OCT-2001
                                                                                                                                                                                                                                                                                         AAI52287;
                                                                                                                                                                                                                                                                                                                       AAI52287 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cttcttctcgccgcaagagccgtgccgctcacttcaaggctccctctgaccagcgccgtg 135
                                                                                                                                                                                                                                                                                                                                                                                                                                                             agatcacctccgtgtaccgtctcaagtacgtcgtccacgttgagcgtgtcacccgtgaca
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                                                                              -2001
                                                                                                                                                                                          microarray;
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                                                                                                                                                                                          human;
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Pred. No. 8.8e-23;
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04-FEB-2000; 2000US-0180312 26-MAY-2000; 2000US-0207456

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RESULT
AAF12354
ID AAF1
XX AAF1
XX AAF1
XX AAF1
XX ASPA
XX SASPA
XW ASPA
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KW Cul.
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Best Local
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03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to single exon nucleic acid probes (SENP). The present sequence is one such probe. The probes are useful for producing a microarray for predicting, measuring and displaying gene expression in samples derived from human placenta. The probes are useful for antenatal diagnosis of human genetic disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human genome-derived single exon analyzing gene expression in huma
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                                    Multiple gene expression; filamentous fungal cell; EST; expressed sequence tag; Fusarium venenatum; Aspergillus niger; Aspergillus oryzae; Trichoderma reesei; identification; recomb culture condition; environmental stress; spore morphogenesis; metabolic pathway engineering; catabolic pathway engineering;
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                                                                                                                Aspergillus
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                                                                                                                                                                                              AAF12354
             Aspergillus
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                                                                                                                                                                                                                                                            ATTCACCCAAGCAAGGTGGTTA 102
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                                                                                                                                                                                                                                                                                                                          cggatacatatcgagagagtgaccagagagaaaaagcgaatggaatgactgtacctgtggga
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                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene
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2000US-0632366.
2000US-0234687.
2000US-0236359.
2000US-0024263.
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              oryzae
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                                                                                                                  EST SEQ
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                                                                                                                                                                                              ВP
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                                                                 recombination;
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engineering. Using ESTs provides several advantages over genomic or random cDNA clones including elimination of redundancy as one spot on an array equals one gene or open reading frame, and organisation of the microarray based on function of the gene products to facilitate analysis of the results. AAF07478 to AAF11247 represents ESTs from Aspergillus priger; AAF11854 to AAF14878 represents ESTs from Aspergillus oxyzae; and AAF14879 to AAF14875 represents ESTs from Trichoderma reesei, which are all specifically claimed in the present invention.
                                                                                                                                                                               are used in the methods for monitoring differential expression of genes in a first filamentous fungal (FF) cell relative to expression of the same genes in one or more second filamentous fungal cells. Monitoring the global expression of genes from FF cells allows the production potential of the microorganisms to be improved. New genes may be discovered, possible functions of unknown open reading frames can be identified and gene copy number variation and stability can be monitored. The expression of genes can be used to study how FF cells adapt to changes in culture conditions, environmental stress, spore monitored, recombination, metabolic or catabolic pathway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (NOVO)
                                                                                                                                                                                                                                                                                                                                                      expression of genes in a first filamentous fungal (FF) cell relative to expression of the same genes in one or more second filamentous fungal cells. The method uses fluorescence-labeled nucleic acids isolated from the FF cells and a substrate of expressed sequence tags (EST). The ESTs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Monitoring differential expression of uses fluorescence-labeled nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a method for monitoring differential
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 88; Page 2058; 3161pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  substrate of expressed sequence tags
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ВP;
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Α;
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             gggcaagatcaccagcgtttaccgtcttaagtgggttgtccacgtcgagcgtgtcgtccg
                                                           aggaaaagttacaacgtgttatcgaaagaaatatcggatacatatcgagagagtgaccag
                                                                                                        tttacctattcgaaaagaagacgaagtaatcatagttcgcggagctttcaagggtagaga
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cgagaagtccaacggccagagcgttcccctcggtatccacccctccaaggtcgtca
                                                                                             tctccccatccgcaaggacgacgaggtcaccattgtccgtggaggccagaagggccgtga
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172; Conser
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                                                                                                                                                                                                                                                       Score 97.6; DB 21 Pred. No. 2.5e-21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides the protein and coding sequences of a number of apoptosis associated proteins from the yeast Saccharomyces cerevisiae and the fungus Candida albicans. These can be used to identify treatments for fungal and yeast infections, for proliferative diseases and for apoptosis related diseases such as autoimmune diseases, ischaemia and neurodegeneration. The present sequence is one of the S. cerevisiae
          1205
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                                                                                                                                                                                                              1085
                                                                                                                                                                                                                                                                                                               1025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yeast and fungal nucleic acids encoding proteins involved in a pathway leading to programmed cell death, useful for treating proliferative disorders, yeast and fungal infections, or for preventing apoptosis in certain diseases .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1331 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            coding sequences of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Fig 1; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nelissen BJM,
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                       agaaggaaaagttacaacgtgttatcgaaagaaatatcggatacatatcgagagagtgac
                                                                                                                                                                                                                           ttctttacctattcgaaaagaagacgaagtaatcatagttcgcggagctttcaagggtag
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                                                                                               agaaggtaagatttcatctgtttacagattgaaatttgccgttcaagttgacaaggttac
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                                                                                                                                                                                                                                                                                                     acgtcgtgttttgctatctgctccattatctaaggagttgagagctcaatacggtatcaa
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)B; AAG70777.
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s; autoimmune disease; ischaemia; neurodegeneration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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Pred. No. 1
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1.2e-20;
hes 127;
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aagtacaacgtgcgatccatgcccatccgaaaggatgatgaagttcaggttgtacgtgga

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                                                                                                                                                                                     mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and secretion
                                                                                                                                                                                                                                                              The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. An ORF has been identified within the sequence. The 5' ESTs were prepared from the sequences or polya+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GEST
                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID 3876; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                                                                                                                                                              diagnostic,
                                                                                                                                                                                                                                                                                                                                                                                                        New nucleic acid that is a 5' expressed sequence tag (5'\ EST) j obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dumas Milne Edwards
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 P-PSDB;
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              156
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                                                  atgaagtattccaaagttgtatcgtcgtctagaagaaagcagaggaaggcatactttcaa 155
atgaagtttaatccctttgtgacttccgaccgaagcaagaatcgcaaaaggcatttcaat 127
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                                                                                        182;
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                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                              forensic,
                                                                                                                                                      B₽;
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                                                                                                                                                    172 A;
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                                                                                                 22.2%;
                                                                                                                                                                                                                                                                                                                                                                                            gene therapy and chromosome mapping
                                                                                                                                                    113 C;
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The present invention describes a method for monitoring differential expression of genes in a first filamentous fungal (FF) cell relative to expression of the same genes in one or more second filamentous fungal (cells. The method uses fluorescence-labeled nucleic acids isolated from the FF cells and a substrate of expressed sequence tags (EST). The ESTS care used in the methods for monitoring differential expression of genes in a first filamentous fungal (FF) cell relative to expression of the same genes in one or more second filamentous fungal cells. Monitoring ct the global expression of genes from FF cells allows the production contential of the microorganisms to be improved. New genes may be considered, possible functions of unknown open reading frames can be identified and gene copy number variation and stability can be compitored. The expression of genes can be used to study how FF cells campitored. The expression of genes can be used to study how FF cells campitored. The expression of genes can be used to study how FF cells campitored. The expression of genes can be used to study how FF cells campitored. The expression of genes can be used to study how FF cells cappite to changes in culture conditions, environmental stress, spore corphogenesis, recombination, metabolic or catabolic pathway cells in a culture condition of redundancy as one spot on an array equals one gene or open reading frame, and organisation of the mitroarrays based on function of the gene products to facilitate can allysis of the results. AAF0148 to AAF1127 represents ESTs from Aspergillus oryzae; and creaming the composition of the gene products to facilitate to an allowantes of the results. AFF1248 to AAF1153 represents ESTs from Aspergillus oryzae; and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Multiple gene expression; filamentous fungal cell; EST; expressed sequence tag; Fusarium venenatum; Aspergillus niger; Aspergillus oryzae; Trichoderma reesei; identification; recombination; culture condition; environmental stress; spore morphogenesis; metabolic pathway engineering; catabolic pathway engineering; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Monitoring differential expression of genes uses fluorescence-labeled nucleic acids isol substrate of expressed sequence tags -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Probe; microarray; human; placenta; antenatal diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAI39236;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAI39236 standard; DNA;
                                                                                                               Human genome-derived single exon nucleic acid analyzing gene expression in human placenta -
                                                                                                                                                                                                                                                                                                                             04-OCT-2000;
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                                                                      Claim
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27-SEP-2000;
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  single exon nucleic acid probes (SENP). probe. The probes are useful for
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PR 23-JUL-1999; 99US-0145224.
PR 27-JUL-1999; 99US-0145218.
PR 27-JUL-1999; 99US-0145918.
PR 27-JUL-1999; 99US-0145919.
PR 28-JUL-1999; 99US-0145919.
PR 28-JUL-1999; 99US-0145919.
PR 02-AUG-1999; 99US-0147303.
PR 03-AUG-1999; 99US-0147303.
PR 06-AUG-1999; 99US-0147303.
PR 06-AUG-1999; 99US-0147303.
PR 07-AUG-1999; 99US-0147303.
PR 11-AUG-1999; 99US-0147303.
PR 11-AUG-1999; 99US-0147416.
PR 11-AUG-1999; 99US-0147416.
PR 11-AUG-1999; 99US-0148317.
PR 11-AUG-1999; 99US-0148317.
PR 11-AUG-1999; 99US-0148364.
PR 12-AUG-1999; 99US-0148364.
PR 13-AUG-1999; 99US-0148364.
PR 23-AUG-1999; 99US-0149368.
PR 23-AUG-1999; 99US-015066.
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28-OCT-1999;
28-OCT-1999;
28-OCT-1999;
29-OCT-1999;
                                                                                                                                                                                                               Multiple gene expression; filamentous fungal cell; EST; expressed sequence tag; Fusarium venenatum; Aspergillus niger; Aspergillus oryzae; Trichoderma reesei; identification; recombination; culture condition; environmental stress; spore morphogenesis; metabolic pathway engineering; catabolic pathway engineering; SS.
                                                                                                                                                                                                                                                                                                   13-MAR-2001
                                                                                                                                                                                                                                                                              Aspergillus niger EST SEQ ID
                                                                                                                                                                                                                                                                                                                        AAF11388
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Monitoring differential expression of genes in filamentous uses fluorescence-labeled nucleic acids isolated from the cubstrate of expressed sequence tags \dot{}
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                                                                                                                                                                                              Aspergillus niger.
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CC same genes in one or more second filamentous fungal (rr) cert retactive to expression of the global expression of genes from FF cells allows the production contential of the microorganisms to be improved. New genes may be cold discovered, possible functions of unknown open reading frames can be continited and gene copy number variation and stability can be compined. The expression of genes can be used to study how FF cells cadapt to changes in culture conditions, environmental stress, spore compined to changes in culture conditions, environmental stress, spore compined to changes in culture conditions, environmental stress, spore compined to change stream of the gene or catabolic or catabolic pathway compined to the compined to the condition of the gene of the condition of the gene products to facilitate of analysis of the results. AAF07478 to AAF11247 represents ESTs from Aspergillus organisation of the canalysis of the results. AAF07478 to AAF11253 represents ESTs from Aspergillus organisation of construm venenatum; AAF11248 to AAF11853 represents ESTs from Aspergillus organisation of construments and construments of the construction of the construments of the construction of the construments of the co
Sequence 629 BP; 153 A; 191 C; 155 G; 129 T; 1 other;
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Qy 밁 QΥ Вb ρ В Ş Вþ δÃ Query Match
Best Local Similarity 54.7
Matches 146; Conservative 310 353 agaaaaagcgaatggaatgactgtacc 379 250 gggcaaggtcaccagcgtctaccgcctgaagtgggccatccacgtcgagcgcatcagccg 309 190 catgcccatccgcaaggacgacgaggtcatggtcgtccgtggcagcaacaagggccgtga 233 130 113 tgtatcgtcgtctagaagaagcagaggaaagcatactttcaagcaccatcttctgtacg 172 70 cgacaagagcaacggccagagcgtccc aggaaaagttacaacgtgttatcgaaagaaatatcggatacatatcgagagagtgaccag 352 tttacctattcgaaaagaagaagtaatcatagttcgcggagctttcaagggtagaga ccgcgttatcctctctctctctcgagcgagctccgcgccaagtacaatgtccgctc 189 18.0%; Score 73.4; DB 21; Length Pred. No. 1.2e-13; 0; Mismatches 121; Indels 336 Length 629; 0; Gaps 249 292 0;

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